

SEQUENCE LISTING

<110> Ajinomoto Co., Inc.

<120> Method for Producing L-Amino Acid

<130>

<160> 24

<210> 1

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yahN gene

<400> 1

ggcgagctcc cagtaaccgg aaataag

27

<210> 2

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yahN gene

<400> 2

cgctctagaa aggaccacgc attacgg

27

<210> 3

<211> 27

<212> DNA

09459573 121399

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yeaS gene

<400> 3

ggcgagctca gattggttag catattc

27

<210> 4

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yeaS gene

<400> 4

cggctctagaa tcagcgaaga atcaggg

27

<210> 5

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yfiK gene

<400> 5

ggcgagctca tggtccgtgt cgggtac

27

<210> 6

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yfiK gene

09459573 121399

<400> 6

ggctctagat agcaagttac taagcgg

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<210> 7

<211> 35

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yggA gene

<400> 7

ctctgaattc tctcttatta gtttttctga ttgcc

35

<210> 8

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yggA gene

<400> 8

cgtgacctgc agcgttctca cagcgcgcta gcctttaa

38

<210> 9

<211> 672

<212> DNA

<213> Escherichia coli

<220>

<221> CDS

<222> (1)..(672)

<400> 9

atg	atg	cag	tta	ggt	cac	tta	ttt	atg	gat	gaa	atc	act	atg	gat	cct	48
Met	Met	Gln	Leu	Val	His	Leu	Phe	Met	Asp	Glu	Ile	Thr	Met	Asp	Pro	
1				5				10					15			
ttg	cat	gcc	ggt	tac	ctg	acc	gta	gga	ctg	ttc	gtg	att	act	ttt	ttt	96
Leu	His	Ala	Val	Tyr	Leu	Thr	Val	Gly	Leu	Phe	Val	Ile	Thr	Phe	Phe	

09459573 121399

20 25 30
 aat ccg gga gcc aat ctc ttt gtg gta gta caa acc agc ctg gct tcc 144
 Asn Pro Gly Ala Asn Leu Phe Val Val Val Gln Thr Ser Leu Ala Ser
 35 40 45
 ggt cga cgc gca ggg gtg ctg acc ggg ctg ggc gtg gcg ctg ggc gat 192
 Gly Arg Arg Ala Gly Val Leu Thr Gly Leu Gly Val Ala Leu Gly Asp
 50 55 60
 gca ttt tat tcc ggg ttg ggt ttg ttt ggt ctt gca acg cta att acg 240
 Ala Phe Tyr Ser Gly Leu Gly Leu Phe Gly Leu Ala Thr Leu Ile Thr
 65 70 75 80
 cag tgt gag gag att ttt tcg ctt atc aga atc gtc ggc ggc gct tat 288
 Gln Cys Glu Glu Ile Phe Ser Leu Ile Arg Ile Val Gly Gly Ala Tyr
 85 90 95
 ctc tta tgg ttt gcg tgg tgc agc atg cgc cgc cag tca aca ccg caa 336
 Leu Leu Trp Phe Ala Trp Cys Ser Met Arg Arg Gln Ser Thr Pro Gln
 100 105 110
 atg agc aca cta caa caa ccg att agc gcc ccc tgg tat gtc ttt ttt 384
 Met Ser Thr Leu Gln Gln Pro Ile Ser Ala Pro Trp Tyr Val Phe Phe
 115 120 125
 cgc cgc gga tta att acc gat ctc tct aac ccg caa acc gtt tta ttt 432
 Arg Arg Gly Leu Ile Thr Asp Leu Ser Asn Pro Gln Thr Val Leu Phe
 130 135 140
 ttt atc agt att ttc tca gta aca tta aat gcc gaa aca cca aca tgg 480
 Phe Ile Ser Ile Phe Ser Val Thr Leu Asn Ala Glu Thr Pro Thr Trp
 145 150 155 160
 gca cgt tta atg gcc tgg gcg ggg att gtg ctc gca tca att atc tgg 528
 Ala Arg Leu Met Ala Trp Ala Gly Ile Val Leu Ala Ser Ile Ile Trp
 165 170 175
 cga gtt ttt ctt agt cag gcg ttt tct ttg ccc gct gtg cgt cgt gct 576
 Arg Val Phe Leu Ser Gln Ala Phe Ser Leu Pro Ala Val Arg Arg Ala
 180 185 190
 tat ggg cgt atg caa cgc gtt gcc agt cgg gtt att ggt gca att att 624
 Tyr Gly Arg Met Gln Arg Val Ala Ser Arg Val Ile Gly Ala Ile Ile
 195 200 205
 ggt gta ttc gcg cta cgc ctg att tac gaa ggg gtg acg cag cgg tga 672
 Gly Val Phe Ala Leu Arg Leu Ile Tyr Glu Gly Val Thr Gln Arg
 210 215 220

<210> 10

<211> 223

<212> PRT

<213> Escherichia coli

0945953 121399

<400> 10

Met Met Gln Leu Val His Leu Phe Met Asp Glu Ile Thr Met Asp Pro
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 Leu His Ala Val Tyr Leu Thr Val Gly Leu Phe Val Ile Thr Phe Phe
 20 25 30
 Asn Pro Gly Ala Asn Leu Phe Val Val Val Gln Thr Ser Leu Ala Ser
 35 40 45
 Gly Arg Arg Ala Gly Val Leu Thr Gly Leu Gly Val Ala Leu Gly Asp
 50 55 60
 Ala Phe Tyr Ser Gly Leu Gly Leu Phe Gly Leu Ala Thr Leu Ile Thr
 65 70 75 80
 Gln Cys Glu Glu Ile Phe Ser Leu Ile Arg Ile Val Gly Gly Ala Tyr
 85 90 95
 Leu Leu Trp Phe Ala Trp Cys Ser Met Arg Arg Gln Ser Thr Pro Gln
 100 105 110
 Met Ser Thr Leu Gln Gln Pro Ile Ser Ala Pro Trp Tyr Val Phe Phe
 115 120 125
 Arg Arg Gly Leu Ile Thr Asp Leu Ser Asn Pro Gln Thr Val Leu Phe
 130 135 140
 Phe Ile Ser Ile Phe Ser Val Thr Leu Asn Ala Glu Thr Pro Thr Trp
 145 150 155 160
 Ala Arg Leu Met Ala Trp Ala Gly Ile Val Leu Ala Ser Ile Ile Trp
 165 170 175
 Arg Val Phe Leu Ser Gln Ala Phe Ser Leu Pro Ala Val Arg Arg Ala
 180 185 190
 Tyr Gly Arg Met Gln Arg Val Ala Ser Arg Val Ile Gly Ala Ile Ile
 195 200 205
 Gly Val Phe Ala Leu Arg Leu Ile Tyr Glu Gly Val Thr Gln Arg
 210 215 220

<210> 11

<211> 639

<212> DNA

<213> Escherichia coli

<220>

<221> CDS

<222> (1)..(639)

<400> 11

gtg ttc gct gaa tac ggg gtt ctg aat tac tgg acc tat ctg gtt ggg 48

09459573-121399

<210> 12
 <211> 212
 <212> PRT
 <213> Escherichia coli

<400> 12

Met Phe Ala Glu Tyr Gly Val Leu Asn Tyr Trp Thr Tyr Leu Val Gly
 1 5 10 15
 Ala Ile Phe Ile Val Leu Val Pro Gly Pro Asn Thr Leu Phe Val Leu
 20 25 30
 Lys Asn Ser Val Ser Ser Gly Met Lys Gly Gly Tyr Leu Ala Ala Cys
 35 40 45
 Gly Val Phe Ile Gly Asp Ala Val Leu Met Phe Leu Ala Trp Ala Gly
 50 55 60
 Val Ala Thr Leu Ile Lys Thr Thr Pro Ile Leu Phe Asn Ile Val Arg
 65 70 75 80
 Tyr Leu Gly Ala Phe Tyr Leu Leu Tyr Leu Gly Ser Lys Ile Leu Tyr
 85 90 95
 Ala Thr Leu Lys Gly Lys Asn Ser Glu Ala Lys Ser Asp Glu Pro Gln
 100 105 110
 Tyr Gly Ala Ile Phe Lys Arg Ala Leu Ile Leu Ser Leu Thr Asn Pro
 115 120 125
 Lys Ala Ile Leu Phe Tyr Val Ser Phe Phe Val Gln Phe Ile Asp Val
 130 135 140
 Asn Ala Pro His Thr Gly Ile Ser Phe Phe Ile Leu Ala Ala Thr Leu
 145 150 155 160
 Glu Leu Val Ser Phe Cys Tyr Leu Ser Phe Leu Ile Ile Ser Gly Ala
 165 170 175
 Phe Val Thr Gln Tyr Ile Arg Thr Lys Lys Lys Leu Ala Lys Val Gly
 180 185 190
 Asn Ser Leu Ile Gly Leu Met Phe Val Gly Phe Ala Ala Arg Leu Ala
 195 200 205
 Thr Leu Gln Ser
 210

<210> 13
 <211> 588
 <212> DNA
 <213> Escherichia coli

<220>
 <221> CDS

06459573.12139

<400> 13

[illegible]

<210> 14
 <211> 195
 <212> PRT
 <213> Escherichia coli

<400> 14
 Met Thr Pro Thr Leu Leu Ser Ala Phe Trp Thr Tyr Thr Leu Ile Thr
 1 5 10 15
 Ala Met Thr Pro Gly Pro Asn Asn Ile Leu Ala Leu Ser Ser Ala Thr
 20 25 30
 Ser His Gly Phe Arg Gln Ser Thr Arg Val Leu Ala Gly Met Ser Leu
 35 40 45
 Gly Phe Leu Ile Val Met Leu Leu Cys Ala Gly Ile Ser Phe Ser Leu
 50 55 60
 Ala Val Ile Asp Pro Ala Ala Val His Leu Leu Ser Trp Ala Gly Ala
 65 70 75 80
 Ala Tyr Ile Val Trp Leu Ala Trp Lys Ile Ala Thr Ser Pro Thr Lys
 85 90 95
 Glu Asp Gly Leu Gln Ala Lys Pro Ile Ser Phe Trp Ala Ser Phe Ala
 100 105 110
 Leu Gln Phe Val Asn Val Lys Ile Ile Leu Tyr Gly Val Thr Ala Leu
 115 120 125
 Ser Thr Phe Val Leu Pro Gln Thr Gln Ala Leu Ser Trp Val Val Gly
 130 135 140
 Val Ser Val Leu Leu Ala Met Ile Gly Thr Phe Gly Asn Val Cys Trp
 145 150 155 160
 Ala Leu Ala Gly His Leu Phe Gln Arg Leu Phe Arg Gln Tyr Gly Arg
 165 170 175
 Gln Leu Asn Ile Val Leu Ala Leu Leu Leu Val Tyr Cys Ala Val Arg
 180 185 190
 Ile Phe Tyr
 195

<210> 15
 <211> 636
 <212> DNA
 <213> Escherichia coli

<220>
 <221> CDS
 <222> (1)..(636)

00459573.121399

<400> 15
 gtg ttt tct tat tac ttt caa ggt ctt gca ctt ggg gcg gct atg atc 48
 Met Phe Ser Tyr Tyr Phe Gln Gly Leu Ala Leu Gly Ala Ala Met Ile
 1 5 10 15
 cta ccg ctc ggt cca caa aat gct ttt gtg atg aat cag ggc ata cgt 96
 Leu Pro Leu Gly Pro Gln Asn Ala Phe Val Met Asn Gln Gly Ile Arg
 20 25 30
 cgt cag tac cac att atg att gcc tta ctt tgt gct atc agc gat ttg 144
 Arg Gln Tyr His Ile Met Ile Ala Leu Leu Cys Ala Ile Ser Asp Leu
 35 40 45
 gtc ctg att tgc gcc ggg att ttt ggt ggc agc gcg tta ttg atg cag 192
 Val Leu Ile Cys Ala Gly Ile Phe Gly Gly Ser Ala Leu Leu Met Gln
 50 55 60
 tcg ccg tgg ttg ctg gcg ctg gtc acc tgg ggc ggc gta gcc ttc ttg 240
 Ser Pro Trp Leu Leu Ala Leu Val Thr Trp Gly Gly Val Ala Phe Leu
 65 70 75 80
 ctg tgg tat ggt ttt ggc gct ttt aaa aca gca atg agc agt aat att 288
 Leu Trp Tyr Gly Phe Gly Ala Phe Lys Thr Ala Met Ser Ser Asn Ile
 85 90 95
 gag tta gcc agc gcc gaa gtc atg aag caa ggc aga tgg aaa att atc 336
 Glu Leu Ala Ser Ala Glu Val Met Lys Gln Gly Arg Trp Lys Ile Ile
 100 105 110
 gcc acc atg ttg gca gtg acc tgg ctg aat ccg cat gtt tac ctg gat 384
 Ala Thr Met Leu Ala Val Thr Trp Leu Asn Pro His Val Tyr Leu Asp
 115 120 125
 act ttt gtt gta ctg ggc agc ctt ggc ggg caa ctt gat gtg gaa cca 432
 Thr Phe Val Val Leu Gly Ser Leu Gly Gly Gln Leu Asp Val Glu Pro
 130 135 140
 aaa cgc tgg ttt gca ctc ggg aca att agc gcc tct ttc ctg tgg ttc 480
 Lys Arg Trp Phe Ala Leu Gly Thr Ile Ser Ala Ser Phe Leu Trp Phe
 145 150 155 160
 ttt ggt ctg gct ctt ctc gca gcc tgg ctg gca ccg cgt ctg cgc acg 528
 Phe Gly Leu Ala Leu Leu Ala Ala Trp Leu Ala Pro Arg Leu Arg Thr
 165 170 175
 gca aaa gca cag cgc att atc aat ctg gtt gtg gga tgt gtt atg tgg 576
 Ala Lys Ala Gln Arg Ile Ile Asn Leu Val Val Gly Cys Val Met Trp
 180 185 190
 ttt att gcc ttg cag ctg gcg aga gac ggt att gct cat gca caa gcc 624
 Phe Ile Ala Leu Gln Leu Ala Arg Asp Gly Ile Ala His Ala Gln Ala
 195 200 205
 ttg ttc agt tag 636

0045953" 12130

Leu Phe Ser
210

<210> 16

<211> 211

<212> PRT

<213> Escherichia coli

<400> 16

Met Phe Ser Tyr Tyr Phe Gln Gly Leu Ala Leu Gly Ala Ala Met Ile
1 5 10 15
Leu Pro Leu Gly Pro Gln Asn Ala Phe Val Met Asn Gln Gly Ile Arg
20 25 30
Arg Gln Tyr His Ile Met Ile Ala Leu Leu Cys Ala Ile Ser Asp Leu
35 40 45
Val Leu Ile Cys Ala Gly Ile Phe Gly Gly Ser Ala Leu Leu Met Gln
50 55 60
Ser Pro Trp Leu Leu Ala Leu Val Thr Trp Gly Gly Val Ala Phe Leu
65 70 75 80
Leu Trp Tyr Gly Phe Gly Ala Phe Lys Thr Ala Met Ser Ser Asn Ile
85 90 95
Glu Leu Ala Ser Ala Glu Val Met Lys Gln Gly Arg Trp Lys Ile Ile
100 105 110
Ala Thr Met Leu Ala Val Thr Trp Leu Asn Pro His Val Tyr Leu Asp
115 120 125
Thr Phe Val Val Leu Gly Ser Leu Gly Gly Gln Leu Asp Val Glu Pro
130 135 140
Lys Arg Trp Phe Ala Leu Gly Thr Ile Ser Ala Ser Phe Leu Trp Phe
145 150 155 160
Phe Gly Leu Ala Leu Leu Ala Ala Trp Leu Ala Pro Arg Leu Arg Thr
165 170 175
Ala Lys Ala Gln Arg Ile Ile Asn Leu Val Val Gly Cys Val Met Trp
180 185 190
Phe Ile Ala Leu Gln Leu Ala Arg Asp Gly Ile Ala His Ala Gln Ala
195 200 205
Leu Phe Ser
210

<210> 17

<211> 20

<212> DNA

<213> Artificial Sequence

09459573.121300

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yahN gene

<400> 17

gtgtggaacc gacgccggat

20

<210> 18

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yahN gene

<400> 18

tgttgatatg tacggggttc gag

23

<210> 19

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yeaS gene

<400> 19

ctttgccaat cccgtctccc

20

<210> 20

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yeaS gene

<400> 20

09459573.121399

gccccatgca taacggaaag

20

<210> 21

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yfiK gene

<400> 21

gaagatcttg taggccgat aagcgc

26

<210> 22

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yfiK gene

<400> 22

tggttttacc aattggccgc

20

<210> 23

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yggA gene

<400> 23

acttctcccg cgagccagtt c

21

<210> 24

<211> 21

<212> DNA

<213> Artificial Sequence

09459573.121399

<220>

<223> Description of Artificial Sequence: primer for amplifying Escherichia coli yggA gene

<400> 24

ggcaagctta gcgcctctgt t

21

0945953.121399